



#4

SEQUENCE LISTING

<110> Autogen Research Pty Ltd

<120> Novel genes and their use in the modulation of obesity, diabetes and energy imbalance

<130> 2309315/TDO

<140> 10/039,050

<141> 2001-12-31

<150> 60/141,441

<151> 1999-06-29

<160> 22

<170> PatentIn Ver. 2.0

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Gln Met Ser Cys Trp Pro Lys Pro Leu Ile Pro Glu Leu Glu Arg Gln
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Arg Cys Thr Val Val Thr Pro Lys Val Phe Arg Val Gly Glu Tyr Glu
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caa gtt aca ttt gaa gcc cac ggt cac act gac cca ttt gat gta acc 246
 Gln Val Thr Phe Glu Ala His Gly His Thr Asp Pro Phe Asp Val Thr
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 atc tct ata aaa agt tac cct gat aaa aat gct aat tac tct tca agc 294
 Ile Ser Ile Lys Ser Tyr Pro Asp Lys Asn Ala Asn Tyr Ser Ser Ser
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 Val Tyr Leu Glu Val Val Ser Lys His Phe Ser Thr Ser Lys Ile Met
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Ala Ser Lys Tyr Lys His Pro Val Leu Lys Lys Cys Cys Tyr Asp Gly
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 Ala Arg Tyr Asn His His Glu Thr Cys Glu Glu Arg Val Ala Arg Val
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Ser Leu Leu Ala Ser Tyr Gly Trp Tyr Val Leu Phe Ser Cys Ile Leu
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Leu Tyr Ile Val Ile Gln Lys Leu Ser Val Arg Leu Arg Val Leu Arg
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cag agg cag ctg gac cag gct gac gct gtt ctg gaa cct gat gct gtt 244
Gln Arg Gln Leu Asp Gln Ala Asp Ala Val Leu Glu Pro Asp Ala Val
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Val Lys Arg Gln Glu Ala Leu Ala Ala Arg Leu Arg Met Gln Glu
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95 100 105

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Glu Glu Lys Arg Arg Gln Lys Ile Glu Met Trp Asp Ser Met Gln
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Gly Pro Gly Pro Ser Thr Ser Ser Val Thr Arg Lys Gly Lys Ser
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Asp Lys Lys Pro Leu Arg Gly Asn Gly Tyr Asn Pro Leu Thr Gly Glu
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Gln Lys Leu Ser Val Arg Leu Arg Val Leu Arg Gln Arg Gln Leu Asp
 50 55 60

Gln Ala Asp Ala Val Leu Glu Pro Asp Ala Val Val Lys Arg Gln Glu
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Ala Leu Ala Ala Ala Arg Leu Arg Met Gln Glu Asp Leu Asn Ala Gln
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Val Glu Lys His Lys Glu Lys Leu Arg Gln Leu Glu Glu Lys Arg
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Arg Gln Lys Ile Glu Met Trp Asp Ser Met Gln Glu Gly Arg Ser Tyr
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Thr Ser Ser Ser Val Thr Arg Lys Gly Lys Ser Asp Lys Lys Pro Leu
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Thr Val Gly Ser Leu Leu Ala Thr Tyr Gly Trp Tyr Ile Val Phe Ser
25 30 35 40

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Cys Ile Leu Leu Tyr Val Val Phe Gln Lys Leu Ser Ala Arg Leu Arg
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gcc ttg agg cag agg cag ctg gac cga gct gcg gct gct gtg gaa cct 246
Ala Leu Arg Gln Arg Gln Leu Asp Arg Ala Ala Ala Val Glu Pro
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Asp Val Val Val Lys Arg Gln Glu Ala Leu Ala Ala Arg Leu Lys
75 80 85

atg caa gaa gaa cta aat gcg caa gtt gaa aag cat aag gaa aaa ctg 342
Met Gln Glu Glu Leu Asn Ala Gln Val Glu Lys His Lys Glu Lys Leu
90 95 100

aaa caa ctt gaa gaa gaa aaa agg aga cag aag att gaa atg tgg gac 390
Lys Gln Leu Glu Glu Lys Arg Arg Gln Lys Ile Glu Met Trp Asp
105 110 115 120

A
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140 145 150

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Ser Asp Arg Lys Pro Leu Arg Gly Gly Tyr Asn Pro Leu Ser Gly
155 160 165

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170 175 180

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Ser Gly Gly
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35 40 45
Gln Lys Leu Ser Ala Arg Leu Arg Ala Leu Arg Gln Arg Gln Leu Asp
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Arg Ala Ala Ala Ala Val Glu Pro Asp Val Val Val Lys Arg Gln Glu
65 70 75 80

Ala Leu Ala Ala Ala Arg Leu Lys Met Gln Glu Glu Leu Asn Ala Gln
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Val Glu Lys His Lys Glu Lys Leu Lys Gln Leu Glu Glu Glu Lys Arg
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Arg Gln Lys Ile Glu Met Trp Asp Ser Met Gln Glu Gly Lys Ser Tyr
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Lys Gly Asn Ala Lys Lys Pro Gln Glu Glu Asp Ser Pro Gly Pro Ser
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Thr Val Phe Leu Leu Val Thr Ser Leu Pro His
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